

st98045.ST25
SEQUENCE LISTING

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MULTON, Marie-Christine
POLARD-HOuset, Valerie

<120> MEKK1-INTERACTING FHA PROTEIN (M1F1)

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<140> 09/744,125

<141> 2001-01-19

<150> PCT/EP99/05142

<151> 1999-07-21

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 Ala Gly Leu Asp Leu Asn Ser Ser Asp Asn Gln Ser Gly Gly Ser Thr
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 Ala Ser Lys Gly Arg Tyr Ile Pro Pro His Leu Arg Asn Arg Glu Ala
 35 40 45
 Thr Arg Gly Phe Tyr Asp Lys Asp Ser Ser Gly Trp Ser Ser Ser Lys
 50 55 60
 Asp Lys Asp Ala Tyr Ser Ser Phe Gly Ser Arg Ser Asp Ser Arg Gly
 65 70 75 80

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Lys	Ser	Ser	Phe	Phe	Ser	Asp	Arg	Gly	Ser	Gly	Ser	Arg	Gly	Arg	Phe	85	90	95
Asp	Asp	Arg	Gly	Arg	Ser	Asp	Tyr	Asp	Gly	Ile	Gly	Ser	Arg	Gly	Asp	100	105	110
Arg	Ser	Gly	Phe	Gly	Lys	Phe	Glu	Arg	Gly	Gly	Asn	Ser	Arg	Trp	Cys	115	120	125
Asp	Lys	Ser	Asp	Glu	Asp	Asp	Trp	Ser	Lys	Pro	Leu	Pro	Pro	Ser	Glu	130	135	140
Arg	Leu	Glu	Gln	Glu	Leu	Phe	Ser	Gly	Gly	Asn	Thr	Gly	Ile	Asn	Phe	145	150	155
Glu	Lys	Tyr	Asp	Asp	Ile	Pro	Val	Glu	Ala	Thr	Gly	Asn	Asn	Cys	Pro	165	170	175
Pro	His	Ile	Glu	Ser	Phe	Ser	Asp	Val	Glu	Met	Gly	Glu	Ile	Ile	Met	180	185	190
Gly	Asn	Ile	Glu	Leu	Thr	Arg	Tyr	Thr	Arg	Pro	Thr	Pro	Val	Gln	Lys	195	200	205
His	Ala	Ile	Pro	Ile	Ile	Lys	Glu	Lys	Arg	Asp	Leu	Met	Ala	Cys	Ala	210	215	220
Gln	Thr	Gly	Ser	Gly	Lys	Thr	Ala	Ala	Phe	Leu	Leu	Pro	Ile	Leu	Ser	225	230	235
Gln	Ile	Tyr	Ser	Asp	Gly	Pro	Gly	Glu	Ala	Leu	Arg	Ala	Met	Lys	Glu	245	250	255
Asn	Gly	Arg	Tyr	Gly	Arg	Arg	Lys	Gln	Tyr	Pro	Ile	Ser	Leu	Val	Leu	260	265	270
Ala	Pro	Thr	Arg	Glu	Leu	Ala	Val	Gln	Ile	Tyr	Glu	Glu	Ala	Arg	Lys	275	280	285
Phe	Ser	Tyr	Arg	Ser	Arg	Val	Arg	Pro	Cys	Val	Val	Tyr	Gly	Gly	Ala	290	295	300
Asp	Ile	Gly	Gln	Gln	Ile	Arg	Asp	Leu	Glu	Arg	Gly	Cys	His	Leu	Leu	305	310	315
Val	Ala	Thr	Pro	Gly	Arg	Leu	Val	Asp	Met	Met	Glu	Arg	Gly	Lys	Ile	325	330	335
Gly	Leu	Asp	Phe	Cys	Lys	Tyr	Leu	Val	Leu	Asp	Glu	Ala	Asp	Arg	Met	340	345	350
Leu	Asp	Met	Gly	Phe	Glu	Pro	Gln	Ile	Arg	Arg	Ile	Val	Glu	Gln	Asp	355	360	365
Thr	Met	Pro	Pro	Lys	Gly	Val	Arg	His	Thr	Met	Met	Phe	Ser	Ala	Thr	370	375	380
Phe	Pro	Lys	Glu	Ile	Gln	Met	Leu	Ala	Arg	Asp	Phe	Leu	Asp	Glu	Tyr	385	390	395
																		400

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Ile Phe Leu Ala Val Gly Arg Val Gly Ser Thr Ser Glu Asn Ile Thr
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Gln Lys Val Val Trp Val Glu Glu Ser Asp Lys Arg Ser Phe Leu Leu
420 425 430

Asp Leu Leu Asn Ala Thr Gly Lys Asp Ser Leu Thr Leu Val Phe Val
435 440 445

Glu Thr Lys Lys Gly Ala Asp Ser Leu Glu Asp Phe Leu Tyr His Glu
450 455 460

Gly Tyr Ala Cys Thr Ser Ile His Gly Asp Arg Ser Gln Arg Asp Arg
465 470 475 480

Glu Glu Ala Leu His Gln Phe Arg Ser Gly Lys Ser Pro Ile Leu Val
485 490 495

Ala Thr Ala Val Ala Ala Arg Gly Leu Asp Ile Ser Asn Val Lys His
500 505 510

Val Ile Asn Phe Asp Leu Pro Ser Asp Ile Glu Glu Tyr Val His Arg
515 520 525

Ile Gly Arg Thr Gly Arg Val Gly Asn Leu Gly Leu Ala Thr Ser Phe
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Phe Asn Glu Arg Asn Ile Asn Ile Thr Lys Asp Leu Leu Asp Leu Leu
545 550 555 560

Val Glu Ala Lys Gln Glu Val Pro Ser Trp Leu Glu Asn Met Ala Tyr
565 570 575

Glu His His Tyr Lys Gly Ser Ser Arg Gly Arg Ser Lys Ser Ser Arg
580 585 590

Phe Ser Gly Gly Phe Gly Ala Arg Asp Tyr Arg Gln Ser Ser Gly Ala
595 600 605

Ser Ser Ser Ser Phe Ser Ser Ser Arg Ala Ser Ser Ser Arg Ser Gly
610 615 620

Gly Gly Gly His Gly Ser Ser Arg Gly Phe Gly Gly Gly Gly Tyr Gly
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18